ATGGCTCTGGGGCCCAACTGTGGCATCCTACTGTTTCTGGCTGTTTCTGGGTGTGGC CATCCCCAGGTTTCAAACTCGGGAAGTCGAATCGTGGGAGGGCATGCTGCCCCAGC AGGCACATGGCCGTGGCAGGCTAGCCTCCGTCTGCACAAGGTGCACGTGTGTGGAG GCTCCCTGCTCAGTCCAGAATGGGTGCTCACAGCAGCCCACTGCTTCTCTGGGTCTG TGAACTCGTCTGATTATCAGGTGCACTTGGGAGAGCTTACGGTCACACTGTCTCCCC ACTTCTCCACTGTAAAACGGATCATCATGTACACTGGCTCTCCAGGACCACCGGGGT GTCCAGCCTGTGTGCCTCCCAGAGGCCTCAGCTGACTTCTACCCTGGGATGCAGTGC TGGGTGACTGGCTGGGGCTATACAGGGGAGGGAGAGCCTCTGAAGCCCCCATACAA CCTTCAGGAGGCCAAAGTCTCTGTGGTGGATGTAAAGACCTGCAGCCAGGCTTACAA TAGTCCCAATGGCAGCCTCATCCAGCCAGACATGCTATGCGCCCGGGGCCCTGGGG ATGCCTGCCAGGATGACTCTGGAGGGCCACTAGTCTGCCAGGTGGCTGGAACCTGG CAGCAGGCCGCGTTGTCAGCTGGGGTGAGGGCTGTGGCCGCCCTGACCGCCCTG GCGTCTATGCCCGGGTTACTGCCTATGTAAACTGGATCCACCACCACATCCCGGAAG CAGGGGGCTCAGGAATGCAAGGGCTTCCCTGGGCTCCTCCTCGCCTCTTC TGGCCAAGCCTCTTCCTGCTGCTGGTCTCTGGAGTCCTGATGGCCAAGTACTGGCTG TGTCTTTCTTAAATAAGTTAGTGTTTATTCAGTTTGCTTTGCCCCTCCCCTCCCCTTAG CTTTGACTTAGGAAGCCAAAGTTTTCTGCATCAGATTATTGCAACATTTAACCTGAATT

MALGPNCGILLFLAVSGCGHPQVSNSGSRIVGGHAAPAGTWPWQ

ASLRLHKVHVCGGSLLSPEWVLTAAHCFSGSVNSSDYQVHLGELTVTLSPHFSTVKRI
IMYTGSPGPPGSSGDIALVQLSSPVALSSQVQPVCLPEASADFYPGMQCWVTGWGYTG
EGEPLKPPYNLQEAKVSVVDVKTCSQAYNSPNGSLIQPDMLCARGPGDACQDDSGGPL
VCQVAGTWQQAGVVSWGEGCGRPDRPGVYARVTAYVNWIHHHIPEAGGSGMQGLPWAP
LLAALFWPSLFLLLVSGVLMAKYWLSSPSHAASEL (SEO ID NO:2)

underlined = deleted in targeting construct

[] = sequence flanking Neo insert in targeting construct

ATGGCTCTGGGGCCCAACTGTGGCATCCTACTGTTTCTGGCTGTTTCTG [GGTGTGGCCA TCCCCAGGTTTCAAACTCGGGAAGTCGAATCGTGGGAGGGCATGCTGCCCCAGCAGGCAC ATGGCCGTGGCAGGCTAGCCTCCGTCTGCACAAGGTGCACGTGT]GTGGAGGCTCCCTGC TCAGTCCAGAATGGGTGCTCACAGCAGCCCACTGCTTCTCTGGGTCTGTGAACTCGTCTG ATTATCAGGTGCACTTGGGAGAGCTTACGGTCACACTGTCTCCCCACTT [CTCCACTGTA AAACGGATCATCATGTACACTGGCTCTCCAGGACCACCGGGGTCCAGTGGGGACATTGCC CTGGTGCAGCTGTCCTCCCGGTGGCCCTTTCCAGCCAGGTCCAGCCTGTGTGCCTCCCA GGGGAGGGAG] AGCCTCTGAAGCCCCCATACAACCTTCAGGAGGCCAAAGTCTCTGTGGT CATGCTATGCGCCCGGGGCCCTGGGGATGCCTGCCAGGATGACTCTGGAGGGCCACTAGT CTGCCAGGTGGCTGGAACCTGGCAGCAGGCCGGCGTTGTCAGCTGGGGTGAGGGCTGTGG CCGCCTGACCGCCTGGCGTCTATGCCCGGGTTACTGCCTATGTAAACTGGATCCACCA CCACATCCCGGAAGCAGGGGCTCAGGAATGCAAGGGCTTCCCTGGGCTCCTCTCCTGGC TGCCCTCTTCTGGCCAAGCCTCTTCCTGCTGCTGGTCTCTGGAGTCCTGATGGCCAAGTA CAAGTGTCTTTCTTAAATAAGTTAGTGTTTATTCAGTTTGCTTTTGCCCCTCCCCTCCCCT TAGCTTTGACTTAGGAAGCCAAAGTTTTCTGCATCAGATTATTGCAACATTTAACCTGAA

Gene Sequence Structure *	164 bp	Sequence Deleted	287 bp	٠
Size of full-length cDNA: 1122 bp				,

FIGURE 2A

: 1

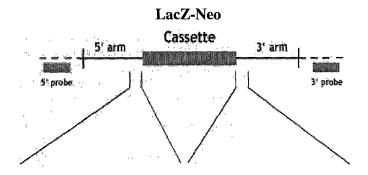
Targeting Vector* (genomic sequence)

Construct Number: 1607

Arm Length: 5': 6 kb 3': 0.7 kb

Targeting Vector
- - - Endogenous Locus

* Not drawn to scale



5'>GGAGTCATGGAGGGCTCCCAG AGAAAGGGCATTGAGCAGAATGCC GGTCTCCAGATTCCCTCACCAACA GTGTCTCCTCTGGATCAGGGTGTG GCCATCCCCAGGTTTCAAACTCGG GAAGTCGAATCGTGGAGGGCATG CTGCCCCAGCAGGCACATGGCCGT GGCAGGCTAGCCTCCGTCTGCACA AGGTGACGTGT<3'

(SEQ ID NO:3)

5'>CTCCACTGTAAAACGGATCAT CATGTACACTGGCTCTCCAGGACC ACCGGGTCCAGTGGGGACATTGC CCTGGTGCAGCTGTCCTCCCGGT GGCCCTTTCCAGCCAGGTCCAGCC TGTGTGCCTCCCAGAGGCCTCAGC TGACTTCTACCCTGGGATGCAGTG CTGGGTGACTGGCTGGGGCTATAC AGGGGAGGGAG<3' (SEQ ID NO:4)

....

necropsy - body weight

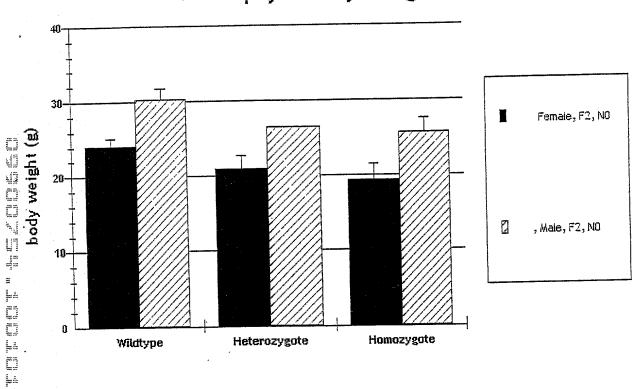


FIGURE 3

necropsy - thymus weight/body weight

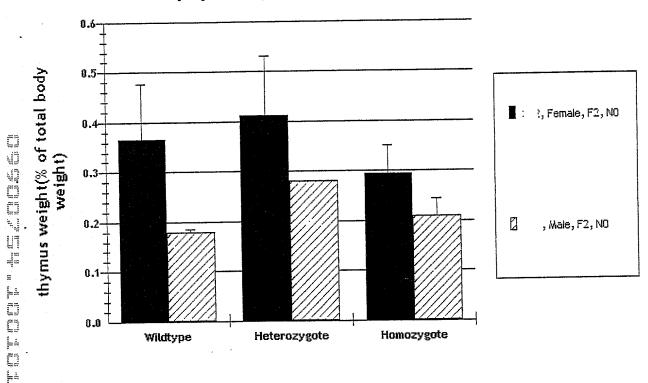


FIGURE 4

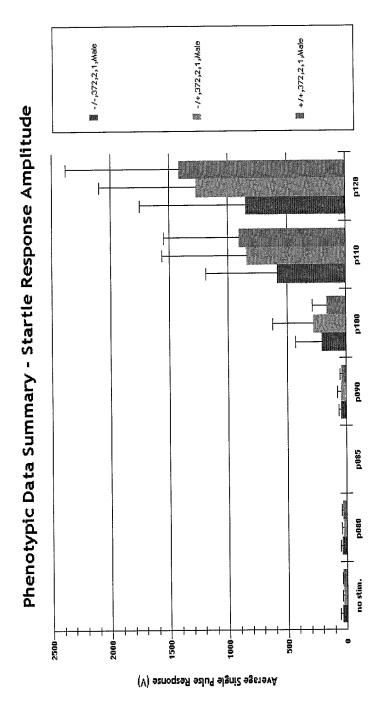


FIGURE 5